

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 09/895,435B
Source: 1FW16
Date Processed by STIC: 12/29/04

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 12/29/2004

PATENT APPLICATION: US/09/895,435B

TIME: 14:51:43

Input Set : A:\29248-17 seq list.ST25.txt

Output Set: N:\CRF4\12292004\I895435B.raw

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3 <110> APPLICANT: DE CLERCQ, BRANTS & PARTNERS c.v.
4     STEWART, A. Francis
5     ZHANG, Youming
6     HALLET, Bernard
8 <120> TITLE OF INVENTION: TYROSINE RECOMBINASE FOR GENETIC ENGINEERING
10 <130> FILE REFERENCE: 29248/17
12 <140> CURRENT APPLICATION NUMBER: US 09/895,435B
13 <141> CURRENT FILING DATE: 2001-06-30
15 <160> NUMBER OF SEQ ID NOS: 10
17 <170> SOFTWARE: PatentIn version 3.2
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 244
21 <212> TYPE: DNA
22 <213> ORGANISM: Bacillus thuringiensis
24 <400> SEQUENCE: 1
25 ggtaccgcca gcatttcgga aaaaaaccac gctaagaaaa tcagagttaa aaaatcagaa      60
27 aatatatcat tattccttga cacatacatg ttcttttttt atacaaaaaa taataacaaca      120
29 caatattaat tgtgttgtat taggtgttat aataaatata aatctagggg tttaacgcaa      180
31 cacaatttat cgataaataa atacttttag acgcaacaca atttatagac gcggaggaaa      240
33 tcac                                                                    244
36 <210> SEQ ID NO: 2
37 <211> LENGTH: 118
38 <212> TYPE: DNA
39 <213> ORGANISM: Bacillus thuringiensis
41 <400> SEQUENCE: 2
42 taataacaaca caatattaat tgtgttgtat taggtgttat aataaatata aatctagggg      60
44 tttaacgcaa cacaatttat cgataaataa atacttttag acgcaacaca atttatag      118
47 <210> SEQ ID NO: 3
48 <211> LENGTH: 32
49 <212> TYPE: DNA
50 <213> ORGANISM: Bacillus thuringiensis
52 <400> SEQUENCE: 3
53 taataacaaca caatattaat tgtgttgtat ta                                                                    32
56 <210> SEQ ID NO: 4
57 <211> LENGTH: 249
58 <212> TYPE: DNA
59 <213> ORGANISM: Bacillus thuringiensis
61 <400> SEQUENCE: 4
62 ggggtaccgc cagcatttcg gaaaaaaacc acgctaagaa aatcagagtt aaaaaatcag      60
64 aaaatatatc attattccaa gacacataca gtgtcttttt ttatacaaaa aataatacaa      120
66 cacaatatta attgtgttgt attaggtgtt ataataaata taaatctagg ggtttaacgc      180
68 aacacaattt atcgataaat aaatactttt agacgcaaca caatttatag acgcggaggaa      240
70 aatcacatg                                                                    249

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73 <210> SEQ ID NO: 5
74 <211> LENGTH: 798
75 <212> TYPE: DNA
76 <213> ORGANISM: Bacillus thuringiensis
78 <400> SEQUENCE: 5
79 atggatgttg caaaacagtt ttcttcttat cttaaacaag agaataaaac cgagaacact    60
81 gttcagggat acacatcagg tattagacag tacataaaat ggtttgaagg ttcctatgac    120
83 agaaaattga caaaattgta cgcacaaaat atcttagagt acattagtta tttaaagaat    180
85 gtcaaaatgt tgaacgccaa gtccattaac cacaagatta gtagccttgc taaatttaat    240
87 gaattttctaa tacagaaagg aagtcaacaa gatcaagtaa ttttattaga tgtaaaaaaag    300
89 tttttacaaa gtgtgttaga ggataataac aaacgtaatt atgcaattgc cactctccta    360
91 gcatatacag gagtacgtat ttcagaggca ttatctatca aaatgaatga cttcaattta    420
93 cagactgggg aatgtattat tcgaagtgga aaaggaggta aacaacgaat tgtattacta    480
95 aatagtaagg tacttagtgc tatcaaagat tatctcatcg atcgaaaaac atacagtaca    540
97 gcatatgaat ctccgtatct ttttattagt aaaaagcgag aaaagctcga ccgtacggtc    600
99 gtcaatcgta tctttaaatc atacaggaat gttattactc cacaccaatt acgacacttc    660
101 ttctgtacga atgcaattca aaaaggattt agcattcatg aagttgcaaa tcaagctggg    720
103 cactctaaca tccatacgac actactttac acaaatccaa accaactgca gctaaaaaat    780
105 aaaatggagc tcttataa                                798

108 <210> SEQ ID NO: 6
109 <211> LENGTH: 265
110 <212> TYPE: PRT
111 <213> ORGANISM: Bacillus thuringiensis
113 <400> SEQUENCE: 6
115 Met Asp Val Ala Lys Gln Phe Ser Ser Tyr Leu Lys Gln Glu Asn Lys
116 1          5          10          15
119 Thr Glu Asn Thr Val Gln Gly Tyr Thr Ser Gly Ile Arg Gln Tyr Ile
120          20          25          30
123 Lys Trp Phe Glu Gly Ser Tyr Asp Arg Lys Leu Thr Lys Leu Tyr Arg
124          35          40          45
127 Gln Asn Ile Leu Glu Tyr Ile Ser Tyr Leu Lys Asn Val Lys Met Leu
128          50          55          60
131 Asn Ala Lys Ser Ile Asn His Lys Ile Ser Ser Leu Ala Lys Phe Asn
132 65          70          75          80
135 Glu Phe Leu Ile Gln Lys Gly Ser Gln Gln Asp Gln Val Ile Leu Leu
136          85          90          95
139 Asp Val Lys Lys Phe Leu Gln Ser Val Leu Glu Asp Asn Asn Lys Arg
140          100         105         110
143 Asn Tyr Ala Ile Ala Thr Leu Leu Ala Tyr Thr Gly Val Arg Ile Ser
144          115         120         125
147 Glu Ala Leu Ser Ile Lys Met Asn Asp Phe Asn Leu Gln Thr Gly Glu
148          130         135         140
151 Cys Ile Ile Arg Ser Gly Lys Gly Gly Lys Gln Arg Ile Val Leu Leu
152 145         150         155         160
155 Asn Ser Lys Val Leu Ser Ala Ile Lys Asp Tyr Leu Ile Asp Arg Lys
156          165         170         175
159 Thr Tyr Ser Thr Ala His Glu Ser Pro Tyr Leu Phe Ile Ser Lys Lys
160          180         185         190
163 Arg Glu Lys Leu Asp Arg Thr Val Val Asn Arg Ile Phe Lys Ser Tyr

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164          195          200          205
167 Arg Asn Val Ile Thr Pro His Gln Leu Arg His Phe Phe Cys Thr Asn
168          210          215          220
171 Ala Ile Gln Lys Gly Phe Ser Ile His Glu Val Ala Asn Gln Ala Gly
172 225          230          235          240
175 His Ser Asn Ile His Thr Thr Leu Leu Tyr Thr Asn Pro Asn Gln Leu
176          245          250          255
179 Gln Leu Lys Asn Lys Met Glu Leu Leu
180          260          265
183 <210> SEQ ID NO: 7
184 <211> LENGTH: 116
185 <212> TYPE: DNA
186 <213> ORGANISM: Bacillus thuringiensis
188 <400> SEQUENCE: 7
189 taataacaaca caatattaat tgtgttgat taggtgttat aataaatata tatctagggg      60
191 ttttaacgcaa cacaatttat cgataaataa atacttttag acgcaacaca atttat      116
194 <210> SEQ ID NO: 8
195 <211> LENGTH: 116
196 <212> TYPE: DNA
197 <213> ORGANISM: Bacillus thuringiensis
199 <400> SEQUENCE: 8
200 taataacaaca caatattaat tgtgttgat taggtgttat aatatatata aatctagggg      60
202 ttttaacgcaa cacaatttat cgataaataa atacttttag acgcaacaca atttat      116
205 <210> SEQ ID NO: 9
206 <211> LENGTH: 116
207 <212> TYPE: DNA
208 <213> ORGANISM: Bacillus thuringiensis
211 <220> FEATURE:
212 <221> NAME/KEY: N_region
213 <222> LOCATION: (33)..(64)
214 <223> OTHER INFORMATION: n = a, t, g, or #c
216 <220> FEATURE:
217 <221> NAME/KEY: misc_feature
218 <222> LOCATION: (33)..(64)
219 <223> OTHER INFORMATION: n is a, c, g, or t
221 <220> FEATURE:
222 <221> NAME/KEY: N_region
223 <222> LOCATION: (81)..(100)
224 <223> OTHER INFORMATION: n = a, t, g, or #c
226 <220> FEATURE:
227 <221> NAME/KEY: misc_feature
228 <222> LOCATION: (81)..(100)
229 <223> OTHER INFORMATION: n is a, c, g, or t
231 <400> SEQUENCE: 9
W--> 232 taataacaaca caatattaat tgtgttgat tannnnnnnnnn nnnnnnnnnnn nnnnnnnnnnn      60
234 nnnnacgcaa cacaatttat nnnnnnnnnnn nnnnnnnnnnn acgcaacaca atttat      116
237 <210> SEQ ID NO: 10
238 <211> LENGTH: 124
239 <212> TYPE: DNA

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240 <213> ORGANISM: Bacillus thuringiensis
 243 <220> FEATURE:
 244 <221> NAME/KEY: N_region
 245 <222> LOCATION: (33)..(74)
 246 <223> OTHER INFORMATION: n = a, t, g, or #c
 248 <220> FEATURE:
 249 <221> NAME/KEY: misc_feature
 250 <222> LOCATION: (33)..(74)
 251 <223> OTHER INFORMATION: n is a, c, g, or t
 253 <400> SEQUENCE: 10

W--> 254 taatacaaca caatattaat tgtgttgat tannnnnnnnn nnnnnnnnnn nnnnnnnnnn 60
 256 nnnnnnnnnn nnnnacgcaa cacaatttac gataaataaa tacttttaga cgcaacacaa 120
 258 tttta 124

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/895,435B

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Input Set : A:\29248-17 seq list.ST25.txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:9; N Pos. ~~33,34,35,36,37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52~~
Seq#:9; N Pos. ~~53,54,55,56,57,58,59,60,61,62,63,64,81,82,83,84,85,86,87,88~~
Seq#:9; N Pos. ~~89,90,91,92,93,94,95,96,97,98,99,100~~
Seq#:10; N Pos. ~~33,34,35,36,37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52~~
Seq#:10; N Pos. ~~53,54,55,56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72~~
Seq#:10; N Pos. ~~73,74~~

VERIFICATION SUMMARY

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Input Set : A:\29248-17 seq list.ST25.txt

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L:232 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0

M:341 Repeated in SeqNo=9

L:254 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0

M:341 Repeated in SeqNo=10